

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 091892,613 D
Source: JFW/C
Date Processed by STIC: 10/02/06

ENTERED



IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/892,613D

DATE: 10/10/2006

TIME: 08:43:49

Input Set : N:\Crf4\10022006\I892613C.raw
 Output Set: N:\CRF4\10102006\I892613D.raw

1 <110> APPLICANT: Leung, Shawn Shui-on
 2 <120> TITLE OF INVENTION: REDUCING IMMUNOGENICITIES OF IMMUNOGLOBULINS BY
 3 FRAMEWORK-PATCHING
 4 <130> FILE REFERENCE: 655
C--> 5 <140> CURRENT APPLICATION NUMBER: US/09/892,613D
 6 <141> CURRENT FILING DATE: 2001-06-27
 7 <160> NUMBER OF SEQ ID NOS: 71
 8 <170> SOFTWARE: PatentIn version 3.3
 10 <210> SEQ ID NO: 1
 11 <211> LENGTH: 369
 12 <212> TYPE: DNA
 13 <213> ORGANISM: Artificial Sequence
 14 <220> FEATURE:
 15 <223> OTHER INFORMATION: FR-patched heavy chain variable region sequence (Full DNA
 16 Sequence) formed by joining the N- and C-terminal (SEQ 3 and 6)
 17 halves at the KpnI site.
 18 <220> FEATURE:
 19 <221> NAME/KEY: V_region
 20 <222> LOCATION: (1)..(369)
 21 <400> SEQUENCE: 1
 22 gaagtgcagc tgctggagtc tgggggaggc tttagtgcagc ctggagggtc cctgaggc 60
 23 tcctgtgcag cctctggatt ctccttcagt atctatgaca tgtcttggtt tcgccaggca 120
 24 ccgggaaagg ggctggagtg ggtcgatac attagtagtg gtgggtgtac cacctactat 180
 25 ccagacactg tgaaggccg attcaccatc tccagagaca atgccaagaa ctccctgtac 240
 26 ctgcaaatga acagtctgag ggtggaggac acagccttat attactgtgc aagacatagt 300
 27 ggctacggta gtagctacgg ggttttgtt gcttactggg gccaaggac tctggtcact 360
 28 gtctcttca 369
 30 <210> SEQ ID NO: 2
 31 <211> LENGTH: 123
 32 <212> TYPE: PRT
 33 <213> ORGANISM: Chimaera sp.
 34 <400> SEQUENCE: 2
 35 Glu Val Gln Leu Leu Glu Ser Gly Gly Leu Val Gln Pro Gly Gly
 36 1 5 10 15
 37 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe Ser Ile Tyr
 38 20 25 30
 39 Asp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 40 35 40 45
 41 Ala Tyr Ile Ser Ser Gly Gly Thr Thr Tyr Tyr Pro Asp Thr Val
 42 50 55 60
 43 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr
 44 65 70 75 80
 45 Leu Gln Met Asn Ser Leu Arg Val Glu Asp Thr Ala Leu Tyr Tyr Cys

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46 85 90 95
 47 Ala Arg His Ser Gly Tyr Gly Ser Ser Tyr Gly Val Leu Phe Ala Tyr
 48 100 105 110
 49 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
 50 115 120
 52 <210> SEQ ID NO: 3
 53 <211> LENGTH: 111
 54 <212> TYPE: DNA
 55 <213> ORGANISM: Artificial Sequence
 56 <220> FEATURE:
 57 <223> OTHER INFORMATION: N-template is a synthetic sense-strand oligonucleotide
 encoding
 58 amino acid 14-50 of the VH region (SEQ ID No. 2). The template
 59 is PCR-amplified by two primers (SEQ ID No. 4 and 5)
 60 <220> FEATURE:
 61 <221> NAME/KEY: V_region
 62 <222> LOCATION: (1)..(111)
 63 <400> SEQUENCE: 3
 64 cctggagggt ccctgaggt ctctgtgca gcctctggat ttccttcag tatstatgac 50
 65 atgtttggg ttccgcaggc accggaaag gggctggagt gggtcgata c TI
 67 <210> SEQ ID NO: 4
 68 <211> LENGTH: 57
 69 <212> TYPE: DNA
 70 <213> ORGANISM: Artificial Sequence
 71 <220> FEATURE:
 72 <223> OTHER INFORMATION: 5' Primer is a synthetic sense-strand oligonucleotide
 encoding
 73 amino acid 1-19 of the VH region (SEQ ID No. 2). The 3' end of
 74 the primer overlaps with the 5'end of the template by 18
 75 nucleotides.
 76 <220> FEATURE:
 77 <221> NAME/KEY: primer_bind
 78 <222> LOCATION: (1)..(57)
 79 <400> SEQUENCE: 4
 80 gaagtgcagc tgctggagtc tgggggaggc tttagtgcagc ctggagggtc cctgagg 57
 82 <210> SEQ ID NO: 5
 83 <211> LENGTH: 48
 84 <212> TYPE: DNA
 85 <213> ORGANISM: Artificial Sequence
 86 <220> FEATURE:
 87 <223> OTHER INFORMATION: 3' Primer is a synthetic anti-sense-strand oligonucleotide
 88 encoding amino acid 43-59 of the VH region(SEQ ID No. 2). The
 89 primer overlaps with the template by 21 nucleotides.
 90 <220> FEATURE:
 91 <221> NAME/KEY: primer_bind
 92 <222> LOCATION: (1)..(48)
 93 <400> SEQUENCE: 5
 94 gttagtgtta ccaccaccac tactaatgta tgcgaccac tccagccc 48
 96 <210> SEQ ID NO: 6
 97 <211> LENGTH: 132
 98 <212> TYPE: DNA

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Input Set : N:\Crf4\10022006\I892613C.raw
 Output Set: N:\CRF4\10102006\I892613D.raw

99 <213> ORGANISM: Artificial Sequence
 100 <220> FEATURE:
 101 <223> OTHER INFORMATION: C-terminal is a synthetic sense-strand oligonucleotide
 encoding
 102 amino acid 68-111 of the VH region (SEQ ID No 2) The template is
 103 PCR-amplified by two primers (SEQ ID No 7 and 8)
 104 <220> FEATURE:
 105 <221> NAME/KEY: V_region
 106 <222> LOCATION: (1)..(132)
 107 <400> SEQUENCE: 6
 108 ttccaccatct ccagagacaa tgccaagaac tccctgtacc tgcaaattgaa cagtttgagg 60
 109 gtggaggaca cagccttata ttactgtgca agacatagtg gctacggtag tagtacggg 120
 110 gtttttttgc ct 132
 112 <210> SEQ ID NO: 7
 113 <211> LENGTH: 60
 114 <212> TYPE: DNA
 115 <213> ORGANISM: Artificial Sequence
 116 <220> FEATURE:
 117 <223> OTHER INFORMATION: 5' Primer is a synthetic sense-strand oligonucleotide
 encoding
 118 amino acid 55-74 of the VH region (SEQ ID No 2). The 3' end of
 119 the primer overlaps with the 5'end of the template by 21
 120 nucleotides.
 121 <220> FEATURE:
 122 <221> NAME/KEY: primer_bind
 123 <222> LOCATION: (1)..(60)
 124 <400> SEQUENCE: 7
 125 ggtggatcca cctactatcc agacactgtg aaggccgat tcaccatctc cagagacaa 60
 127 <210> SEQ ID NO: 8
 128 <211> LENGTH: 57
 129 <212> TYPE: DNA
 130 <213> ORGANISM: Artificial Sequence
 131 <220> FEATURE:
 132 <223> OTHER INFORMATION: 3' Primer is a synthetic anti-sense-strand oligonucleotide
 133 encoding amino acid 105-123 of the VH region (SEQ ID No 2). The
 134 primer and the template overlaps by 21 nucleotides.
 135 <220> FEATURE:
 136 <221> NAME/KEY: primer_bind
 137 <222> LOCATION: (1)..(57)
 138 <400> SEQUENCE: 8
 139 tgaagagaca gtgaccagag tcccttggcc ccagtaagca aacaaaaccc cgttagct 57
 141 <210> SEQ ID NO: 9
 142 <211> LENGTH: 321
 143 <212> TYPE: DNA
 144 <213> ORGANISM: Artificial Sequence
 145 <220> FEATURE:
 146 <223> OTHER INFORMATION: FR-patched light chain variable region sequence formed by
 joining
 147 the N- and C- terminal (SEQ 11 and 14) halves at the KpEI site.
 148 <220> FEATURE:
 149 <221> NAME/KEY: V_region
 150 <222> LOCATION: (1)..(321)

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Output Set: N:\CRF4\10102006\I892613D.raw

151 <400> SEQUENCE: 9
 152 gatatccaga tgaccaggc tccatcctcc ctgtctgcct ctgtgggaga cagagtcacc 60
 153 attagttgca gggcaagtca ggacatttagc aattatttaa actggtatca gcagaaacca 120
 154 ggtaaggctc cgaaactcct gatctactac actagtatat tacactcagg agtcccatca 180
 155 aggttcagtg gcagtgggtc tggAACAGAA ttactctca ccattagctc cctgcagcca 240
 156 gaagattttg ccacttactt ttgccaacag ggtaatacgc ttccgtggac gttcggtgga 300
 157 ggcaccaagg tggAAatcaa a 321

159 <210> SEQ ID NO: 10
 160 <211> LENGTH: 107
 161 <212> TYPE: PRT
 162 <213> ORGANISM: Chimaera sp.
 163 <400> SEQUENCE: 10
 164 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
 165 1 5 10 15
 166 Asp Arg Val Thr Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Asn Tyr
 167 20 25 30
 168 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
 169 35 40 45
 170 Tyr Tyr Thr Ser Ile Leu His Ser Gly Val Pro Ser Arg Phe Ser Gly
 171 50 55 60
 172 Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
 173 65 70 75 80
 174 Glu Asp Phe Ala Thr Tyr Phe Cys Gln Gln Gly Asn Thr Leu Pro Trp
 175 85 90 95
 176 Thr Phe Gly Gly Thr Lys Val Glu Ile Lys
 177 100 105

179 <210> SEQ ID NO: 11
 180 <211> LENGTH: 108
 181 <212> TYPE: DNA
 182 <213> ORGANISM: Artificial Sequence
 183 <220> FEATURE:
 184 <223> OTHER INFORMATION: N-template is a synthetic sense-strand oligonucleotide
 encoding
 185 amino acid 11-46 of the VL region (SEQ ID No. 10). The template
 186 is PCR-amplified by two primers (SEQ ID No. 12 and 13)
 187 <220> FEATURE:
 188 <221> NAME/KEY: V_region
 189 <222> LOCATION: (1)..(108)
 190 <400> SEQUENCE: 11
 191 ctgtctgcct ctgtgggaga cagagtcacc attagttgca gggcaagtca ggacatttagc 60
 192 aattatttaa actggtatca gcagaaacca ggtaaggctc cgaaactc 108
 194 <210> SEQ ID NO: 12
 195 <211> LENGTH: 51
 196 <212> TYPE: DNA
 197 <213> ORGANISM: Artificial Sequence
 198 <220> FEATURE:
 199 <223> OTHER INFORMATION: 5' Primer is a synthetic sense-strand oligonucleotide
 encoding
 200 amino acid 1-17 of the VH region (SEQ ID No 10). The 3' end of
 201 the primer overlaps with the 5'end of the template by 21
 202 nucleotides.

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 PATENT APPLICATION: US/09/892,613D TIME: 08:43:49

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 Output Set: N:\CRF4\10102006\I892613D.raw

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203 <220> FEATURE:
204 <221> NAME/KEY: primer_bind
205 <222> LOCATION: (1)..(51)
206 <400> SEQUENCE: 12
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209 <210> SEQ ID NO: 13
210 <211> LENGTH: 40
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: 3' Primer is a synthetic anti-sense-strand oligonucleotide
215 encoding amino acid 40-53. The primer and the template overlaps
216 by 18 nucleotides.
217 <220> FEATURE:
218 <221> NAME/KEY: primer_bind
219 <222> LOCATION: (1)..(40)
220 <400> SEQUENCE: 13
221 atatactagt gtagtagatc aggagttcg gaggcattacc      40
223 <210> SEQ ID NO: 14
224 <211> LENGTH: 120
225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: C-terminal is a synthetic sense-strand oligonucleotide
encoding
229 amino acid 59-98 of the VH region (SEQ ID No 10) The template is
230 PCR-amplified by tow primers (SEQ ID No 15 and 16)
231 <220> FEATURE:
232 <221> NAME/KEY: V_region
233 <222> LOCATION: (1)..(120)
234 <400> SEQUENCE: 14
235 ccatcaaggt tcagtggcag tgggtctgga acagaattta ctctcaccat tagctccctg      60
236 cagccagaag attttgccac ttactttgc caacagggtt atacgcttcc gtggacgttc      120
238 <210> SEQ ID NO: 15
239 <211> LENGTH: 49
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence
242 <220> FEATURE:
243 <223> OTHER INFORMATION: 5' Primer is a synthetic sense-strand oligonucleotide
encoding
244 amino acid 50-65 of the VH region (SEQ ID No. 10). The 3' end of
245 the primer overlaps with the 5'end of the template by 21
246 nucleotides
247 <220> FEATURE:
248 <221> NAME/KEY: primer_bind
249 <222> LOCATION: (1)..(49)
250 <400> SEQUENCE: 15
251 ctacactagt atattacact caggagtccc atcaaggttc agtggcagt      49
253 <210> SEQ ID NO: 16
254 <211> LENGTH: 48
255 <212> TYPE: DNA

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/10/2006
PATENT APPLICATION: US/09/892,613D TIME: 08:43:50

Input Set : N:\Crf4\10022006\I892613C.raw
Output Set: N:\CRF4\10102006\I892613D.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 15
Seq#:3; Line(s) 57
Seq#:4; Line(s) 72
Seq#:5; Line(s) 87
Seq#:6; Line(s) 101
Seq#:7; Line(s) 117
Seq#:8; Line(s) 132
Seq#:9; Line(s) 146
Seq#:11; Line(s) 184
Seq#:12; Line(s) 199
Seq#:13; Line(s) 214
Seq#:14; Line(s) 228
Seq#:15; Line(s) 243
Seq#:16; Line(s) 258
Seq#:17; Line(s) 272
Seq#:19; Line(s) 314
Seq#:20; Line(s) 329
Seq#:21; Line(s) 344
Seq#:22; Line(s) 358
Seq#:23; Line(s) 374
Seq#:24; Line(s) 390
Seq#:25; Line(s) 404
Seq#:27; Line(s) 443
Seq#:28; Line(s) 459
Seq#:29; Line(s) 474
Seq#:30; Line(s) 488
Seq#:31; Line(s) 503
Seq#:32; Line(s) 518

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/892,613D

DATE: 10/10/2006

TIME: 08:43:50

Input Set : N:\Crf4\10022006\I892613C.raw
Output Set: N:\CRF4\10102006\I892613D.raw

L:5 M:270 C: Current Application Number differs, Wrong Format